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# **Electronic Supplementary Information**

### For

## Strategically Designed Biomodel: Engineering C3-C4 Cleavage of D-Fructose

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Scheme S1. Synthesis of compound 7.





Scheme S3



**Figure S1**. Crystal coordinates of aldolase in association with fructose-bisphosphate (pdb ID 1umg)<sup>6e</sup>. All amino acids have *S*-configuration at  $C_{\alpha}$ . C4-OH of fructose is close to (2.92 Å) phenolic group of Tyr. Hs' are omitted for clarity.

LC-MS of compounds 4-7. Procedure for recording LC-MS is mentioned in the General note.







Figure S3. LC-MS of compound 4b.



Figure S4. LC-MS of compound 4c.







Figure S6. LC-MS of compound 5a.



Figure S7. LC-MS of compound 5b.



Figure S8. LC-MS of compound 5c.



Figure S9. LC-MS of compound 5d.



Figure S10. LC-MS of compound 6a.



Figure S11. LC-MS of compound 6b.



Figure S12. LC-MS of compound 6c.



Figure S13. LC-MS of compound 6d.



Figure S14. LC-MS of compound 7.

# NMR Spectra



Figure S15. <sup>1</sup>H NMR spectrum of compound 11.



Figure S16. <sup>13</sup>C NMR spectrum of compound 11.



Figure S17. DEPT-135 NMR spectrum of compound 11.



Figure S18. <sup>1</sup>H NMR spectrum of compound 12.



Figure S19. <sup>13</sup>C NMR spectrum of compound 12.



Figure S20. <sup>1</sup>H NMR spectrum of compound 13.



Figure S21. <sup>13</sup>C NMR spectrum of compound 13.



Figure S22. DEPT-135 NMR spectrum of compound 13.



Figure S23. <sup>1</sup>H NMR spectrum of compound 14.



Figure S24. <sup>13</sup>C NMR spectrum of compound 14.



Figure S26. <sup>13</sup>C NMR spectrum of compound 15.



Figure S27. DEPT-135 NMR spectrum of compound 15.



Figure S29. <sup>13</sup>C NMR spectrum of compound 16.



Figure S30. DEPT-135 NMR spectrum of compound 16.



Figure S31. <sup>1</sup>H NMR spectrum of compound 18.



Figure S32. <sup>1</sup>H NMR spectrum of compound 19.



Figure S33. <sup>13</sup>C NMR spectrum of compound 19.



Figure S34. DEPT-135 NMR spectrum of compound 19.



Figure S35. <sup>1</sup>H NMR spectrum of compound 6.





Figure S37. <sup>1</sup>H NMR spectrum of compound 4.



Figure S38. <sup>13</sup>C NMR spectrum of compound 4.



Figure S39. DEPT-135 NMR spectrum of compound 4.



Figure S40. HSQC NMR spectrum of compound 4.



**Figure S41.** <sup>1</sup>H-<sup>1</sup>H COSY NMR spectrum of compound **4**.



Figure S42. <sup>1</sup>H-<sup>1</sup>H ROESY NMR spectrum of compound 4.



**Figure S43.** <sup>1</sup>H-<sup>1</sup>H TOCSY NMR spectrum of compound **4**.



Figure S44. IR spectrum of compound 4.



Figure S45. <sup>1</sup>H NMR spectrum of compound 5.



Figure S46. <sup>13</sup>C NMR spectrum of compound 5.



Figure S47. DEPT-135 NMR spectrum of compound 5.



Figure S48. HSQC NMR spectrum of compound 5.



Figure S49. IR spectrum of compound 5.



Figure S50. <sup>1</sup>H NMR spectrum of compound 7.



Figure S52. DEPT-135 NMR spectrum of compound 7.





Figure S53. Mass spectrum of compound 11 (calcd *m/z* 417.1444, [M+H]<sup>+</sup>).



Figure S54. Mass spectrum of compound 12 (calcd *m/z* 403.1288, [M+H]<sup>+</sup>).



Figure S55. Mass spectrum of compound 13 (calcd *m/z* 474.1659, [M+H]<sup>+</sup>).



Figure S56. Mass spectrum of compound 14 (calcd *m/z* 460.1503, [M+H]<sup>+</sup>).



Figure S57. Mass spectrum of compound 15 (calcd *m/z* 736.2977, [M+H]<sup>+</sup>).



Figure S58. Mass spectrum of compound 16 (calcd *m/z* 701.2581, [M+H]<sup>+</sup>).



Figure S59. Mass spectrum of compound 4 (calcd *m/z* 545.2394, [M+H]<sup>+</sup>).



Figure S60. Mass spectrum of compound 5 (calcd m/z 502.2609,  $[M+H]^+$ ).



Figure S61. Mass spectrum of compound 17 (calcd *m/z* 553.1714, [M+Na]<sup>+</sup>).

![](_page_25_Figure_0.jpeg)

Figure S62. Mass spectrum of compound 18 (calcd *m/z* 517.1717, [M+H]<sup>+</sup>).

![](_page_25_Figure_2.jpeg)

Figure S63. Mass spectrum of compound 19 (calcd *m/z* 793.3191 [M+H]<sup>+</sup>).

![](_page_25_Figure_4.jpeg)

Figure S64. Mass spectrum of compound 6 (calcd m/z 659.2823 [M+H]<sup>+</sup>).

![](_page_26_Figure_0.jpeg)

Figure S65. Mass spectrum of compound 20 (calcd *m/z*, 516.2129 [M+H]<sup>+</sup>).

![](_page_26_Figure_2.jpeg)

Figure S66. Mass spectrum of compound 21 (calcd *m/z*, 502.1972 [M+H]<sup>+</sup>).

![](_page_26_Figure_4.jpeg)

Figure S67. Mass spectrum of compound 22 (calcd *m/z*, 573.2343 [M+H]<sup>+</sup>).

![](_page_27_Figure_0.jpeg)

Figure S68. Mass spectrum of compound 23 (calcd m/z, 559.2187 [M+H]<sup>+</sup>).

![](_page_27_Figure_2.jpeg)

Figure S69. Mass spectrum of compound 24 (calcd *m/z* 736.2977, [M+H]<sup>+</sup>).

![](_page_27_Figure_4.jpeg)

![](_page_27_Figure_5.jpeg)

![](_page_28_Figure_0.jpeg)

Figure S72. Mass spectrum of compound 26 (calcd m/z 616.2766,  $[M+H]^+$ ).

![](_page_28_Figure_2.jpeg)

Figure S73. Mass spectrum of compound 27 (calcd *m/z* 586.2660, [M+H]<sup>+</sup>).

![](_page_29_Figure_0.jpeg)

Figure S74. Mass spectrum of reaction mixture of compound 4 and D-fructose. Peak at m/z 707.2925 corresponds to mass of Schiff base ions (calcd m/z 707.2923, [M]<sup>+</sup>). Peaks at m/z 181.0703 and 545.2391 correspond to m/z of fructose and compound 4 ions, respectively.

![](_page_29_Figure_2.jpeg)

**Figure S75**. Mass spectrum of reaction mixture of compound **6** and D-fructose. Peak at m/z 821.3350 corresponds to mass of Schiff base ions (calcd m/z 821.3352, [M]<sup>+</sup>). Peaks at m/z 181.0712 and 659.2824 correspond to m/z of fructose and compound **6** ions, respectively.

![](_page_29_Figure_4.jpeg)

**Figure S76**. HRMS of the reaction mixture of compound **5d** and L-fructose after 60 min of stirring at 25 - 27 °C.

![](_page_30_Figure_0.jpeg)

Figure S77. Mass spectrum of the reaction mixture of compound 26 (Scheme S5) and D-fructose. Peak at m/z 778.3292 corresponds to mass of Schiff base ions (calcd m/z 778.3294,  $[M]^+$ ). Peaks at m/z 181.0710 and 616.2760 correspond to m/z of fructose and compound 26 ions (calcd m/z 616.2766,  $[M+H]^+$ ), respectively.

![](_page_30_Figure_2.jpeg)

Figure S78. Mass spectrum of the reaction mixture of compound 27 (Scheme S6) and D-fructose. Peak at m/z 748.3194 corresponds to mass of Schiff base ions (calcd m/z 748.3188, [M]<sup>+</sup>). Peaks at m/z 181.0708 and 586.2662 correspond to m/z of fructose and compound 27 ions (calcd m/z 586.2660, [M+H]<sup>+</sup>), respectively.

#### Procedure for Energy minimization of compounds<sup>2</sup>

Compounds were built in the workspace of ArgusLab 4.0.1 and energy minimized using PM3 basis set. For each compound, 20,000 iterations and maximum 20,000 steps were recorded to get the optimized geometry of the molecule.

![](_page_31_Figure_0.jpeg)

Figure S79. Energy minimized structures of Schiff bases obtained from (A) compound 5a, (B) compound 5b and (C) compound 5c.

![](_page_31_Figure_2.jpeg)

**Figure S80.** Energy minimized geometry of Schiff bases: A) Schiff base **8** showing 2.79 Å distance between phenolic O and C4-OH of fructose moiety, B) Schiff base corresponding to compound **4d**. Distance between phenolic O and C4-OH is 5.57 Å, C) Schiff base corresponding to compound **6d**. Distance between phenolic O and C4-OH is 8.33Å. In all the three Schiff bases, the asymmetric carbons are marked as yellow and each has S-configuration.

Energy minimized geometries of Schiff bases obtained from compounds **5d**, **4d** and **6d**, respectively supported the preferred cleavage of fructose by compound **5d**. The phenolic O in Schiff base **8** showed the proximity to C4-OH by 2.79 Å while in the case of Schiff bases of

compound **4d** and **5d**, this distance was 5.57 and 8.33 Å, respectively. In contrast to Schiff base **8**, its analogues obtained from compounds **5a**, **5b** and **5c**, carrying respectively *R*, *R*; *R*, *S* and *S*, *R* configurations at  $C_{\alpha}$  of tyr and lys have phenolic O far away from C4-OH of fructose (Figure S78).

![](_page_32_Figure_1.jpeg)

**Figure S81.** (A) LC chromatogram of thick oil obtained after work up of reaction mixture of compound **5d** and D-fructose. LC chromatogram of commercial samples of: (B) D-glyceraldehyde, (C) Dihydroxyacetone and (D) L-glyceraldehyde.

# Kinetics data for cleavage of D-fructose in presence of aldolase and compound 5d.

For aldolase catalyzed reaction. Reaction mixture (2 ml) was containing fructose-1,6bisphosphate (10<sup>-5</sup> M), NADH (0.5  $\mu$ M), 125  $\mu$ g mixture of glycerol-1-phosphate dehydrogenase and isomerase, 500  $\mu$ M Tris-HCl buffer (pH 7.4) and 50 units of aldolase 33 enzyme. The absorbance change at 340 nm was observed in UV spectrum with varying fructose-1,6-biphosphate conc  $(10^{-5} \text{ M} - 10^{-4} \text{ M})$  and time (from 1 min – 10 min) to obtain K<sub>m</sub> by the graph, 1/rate vs 1/conc.

![](_page_33_Figure_1.jpeg)

**Figure S82**. Change in UV spectrum of reaction mixture containing aldolase (80 units), fructose-1,6-bisphosphate ( $10^{-5}$  M), NADH ( $10^{-5}$  M) in tris-HCl buffer with increase in concentration of fructose-1,6-bisphosphate ( $10^{-5} - 10^{-4}$  M).

For compound 5d catalyzed reaction: formation of Schiff base 8. The reaction mixture was containing compound 5d ( $10^{-5}$  M) and D-Fructose ( $10^{-5}$  M) in 2 ml DMSO:H<sub>2</sub>O (1:9) at pH 6.0-6.5. pH of the reaction mixture was recorded at every step and maintained at 6.0-6.5 using 0.1 N HCl/0.1 N NaOH. Keeping the concentration of compound 5d constant, the concentration of D-fructose was stepwise increased from  $10^{-5}$  M to  $10^{-4}$  M and the corresponding change in UV-vis spectrum at 400 nm was noted. The band at 400 nm (due to compound 5d) underwent change as the formation of Schiff takes place (Table S1).

![](_page_33_Figure_4.jpeg)

Figure S83. (A) UV-vis spectrum of compound 5d ( $10^{-5}$  M, DMSO:H<sub>2</sub>O, 1:9 v/v). (B) Change in UV-vis spectrum of compound 5d at  $10^{-5}$  M (DMSO:H<sub>2</sub>O, 1:9) on stepwise addition of D-fructose (0 - 10 equiv,  $10^{-5}$  M  $- 10^{-4}$  M).

Michaelis –Menton constant (K<sub>m</sub>) was calculated from the Lineweaver-Burk equation of enzyme kinetics:

 $1/V = K_m/V_{max} . 1/[S] + 1/V_{max}$ 

V- Reaction rate (change in absorbance per unit time), K<sub>m</sub>- Michaelis-Menton constant,

 $V_{max}$ - Maximum reaction rate (change in absorbance at maximum substrate concentration per unit time), [S]- Substrate concentration.

 $K_{cat} = V_{max} (sec^{-1})/[M]$ , [M] is concentration of compound 5d at  $V_{max}$ .

Table S1. Kinetics data for the formation of Schiff base 8.

Conc of fructose (µM)	Time interval (min)	V <sub>0</sub> (sec <sup>-1</sup> ) (change in absorbance at 400 nm/time)	$ \begin{array}{c} V_0/E_t (sec^{-1}) \\ E_t - conc of \\ compound \\ \textbf{5d} (10^{-5} M) \end{array} $
10	1	6.66 x 10 <sup>-5</sup>	6.66
20	1	1.08 x 10 <sup>-4</sup>	10.8
30	1	1.35 x 10 <sup>-4</sup>	13.5
40	1	1.5 x 10 <sup>-4</sup>	15.0
50	1	1.75 x 10 <sup>-4</sup>	17.5
60	1	1.86 x 10 <sup>-4</sup>	18.6
70	1	1.9 x 10 <sup>-4</sup>	19.0
80	1	1.95 x 10 <sup>-4</sup>	19.5
90	1	2.01 x 10 <sup>-4</sup>	20.1
100	1	2.08 x 10 <sup>-4</sup> (V <sub>max</sub> )	20.8 (K <sub>cat</sub> )

Table S2. Catalytic efficiency of biomodels for the retroaldol reaction.

S.	Reference	Modular	Substrate	K <sub>m</sub>	K <sub>cat (min</sub> <sup>-1</sup> )
No.		assembly		(µM)	
1	Angew. Chem. Int.	Peptide	OH O	5000	0.13
	Ed. 2009, 48, 922-				
	925				
2	Chem. Commun.	Peptide		1800	2.1 x 10 <sup>-4</sup>
	<b>2001,</b> 769-770		CCH3		

			O OH 	900	4.1 x 10 <sup>-4</sup>
			OCH3		
3	J. Am. Chem. Soc.	Peptide		8	2.3 x 10 <sup>-4</sup>
	<b>2002</b> , 124, 3510-		~ 0.~ 0.~ 0.~ 0		
	3511	Peptide		130	2.0 x 10 <sup>-4</sup>
4	Proc. Natl. Acad. Sci	Aldolase	O OH	25	5.0
	<i>USA</i> <b>1998</b> , 95,	antibody	NMe <sub>2</sub>		
	15351-15355		ОН О	14	1.0
			MeO		
			OH O	150	3.3
			MeO		
				37	0.15
			MeO OH O		
				1	0.3
5	I Mal Dial 2014	Designed	Me <sub>2</sub> N <sup>2</sup> V OH O	2.4	2 2 10-6
5	J. MOI. BIOI. 2014,	Designed		34	3.2 X 10 °
	420, 250-271	Designed	MeO <sup>2</sup> V V OH O	1740	1.5 - 10-5
		retroaldolase		1/40	1.5 X 10 <sup>-5</sup>
		Designed	MeO <sup>2</sup> V V OH O	1570	$1.2 \times 10^{-5}$
		retroaldolase		1370	1.5 X 10 <sup>-</sup>
		Designed	OH O	473	$2.7 \times 10^{-5}$
		retroaldolase	Meo	175	2.7 A 10
		Designed	OH O	880	3.6 x 10 <sup>-6</sup>
		retroaldolase	MeO		
6	present experiments	Aldolase	Fructose-1,6-	16	1620
			bisphosphate		
7	Present	Tripeptide	D-fructose	30	1248
	contribution:	appended			
	Compound 5d	acridine			

## **References:**

- B. S. Furniss, A. J. Hannaford, P. W. G. Smith, A. R. Tatchell in *Vogel's Text Book of Practical Organic Chemistry*. 5<sup>th</sup> Edition. 1989, Addison Wesley Longman Ltd. England. pp761.
- Energy minimization was performed using PM3 basis set of software package ArgusLab4.0.1. ArgusLab 4.0.1, M. A. Thompson, Planaria Software LLC, Seattle, WA 98155.